

Resource Summary Report

Generated by [RRID](#) on Sep 10, 2026

PhenoMan

RRID:SCR_005249

Type: Tool

Proper Citation

PhenoMan (RRID:SCR_005249)

Resource Information

URL: <https://code.google.com/p/phenoman/>

Proper Citation: PhenoMan (RRID:SCR_005249)

Description: An interactive software program that integrates phenotypic data exploration, selection, management and quality control using a unified platform for association studies of rare and common variants.

Abbreviations: PhenoMan

Synonyms: phenoman - Phenotypic data exploration selection management and quality control for association studies of rare and common variants

Resource Type: software resource

Keywords: phenotype, quality control, statistical genetics, association study, python, bio.tools

Availability: GNU General Public License, v3

Resource Name: PhenoMan

Resource ID: SCR_005249

Alternate IDs: biotools:phenoman, OMICS_00301

Alternate URLs: <https://bio.tools/phenoman>

Record Creation Time: 20220129T080229+0000

Record Last Update: 20260905T132532+0000

Ratings and Alerts

No rating or validation information has been found for PhenoMan.

No alerts have been found for PhenoMan.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Uciteli A, et al. (2020) Ontological representation, classification and data-driven computing of phenotypes. Journal of biomedical semantics, 11(1), 15.